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RAW SEQUENCE LISTING

DATE: 08/26/2004

PATENT APPLICATION: US/10/817,525

TIME: 11:47:58

Input Set : A:\099-US1.ST25.txt

Output Set: N:\CRF4\08262004\J817525.raw

3 <110> APPLICANT: Quark Biotech, Inc.
 4 Byk, Tamara
 5 Chajut, Ayelet
 7 <120> TITLE OF INVENTION: sFRP1 and Uses Thereof
 9 <130> FILE REFERENCE: 69222-A; 099/US1
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/817,525
 C--> 11 <141> CURRENT FILING DATE: 2004-04-01
 11 <160> NUMBER OF SEQ ID NOS: 2
 13 <170> SOFTWARE: PatentIn version 3.2
 15 <210> SEQ ID NO: 1
 16 <211> LENGTH: 4469
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 30 accgcaggcc gagggccgcc actggccggg gggaccgggc agcagcttgc ggccgcggag 180
 32 ccgggcaacg ctggggactg cgccttttgt ccccgagggt ccctggaagt ttgcggcagg 240
 34 acgcgcgcgg ggaggcggcg gaggcagccc cgacgtcgcg gagaacaggg cgcagagccg 300
 36 gc atg ggc atc ggg cgc agc gag ggg ggc cgc cgc ggg gcc ctg ggc 347
 37 Met Gly Ile Gly Arg Ser Glu Gly Gly Arg Arg Gly Ala Leu Gly
 38 1 5 10 15
 40 gtg ctg ctg gcg ctg ggc gcg gcg ctt ctg gcc gtg ggc tcg gcc agc 395
 41 Val Leu Leu Ala Leu Gly Ala Ala Leu Leu Ala Val Gly Ser Ala Ser
 42 20 25 30
 44 gag tac gac tac gtg agc ttc cag tcg gac atc ggc ccg tac cag agc 443
 45 Glu Tyr Asp Tyr Val Ser Phe Gln Ser Asp Ile Gly Pro Tyr Gln Ser
 46 35 40 45
 48 ggg cgc ttc tac acc aag cca cct cag tgc gtg gac atc ccc gcg gac 491
 49 Gly Arg Phe Tyr Thr Lys Pro Pro Gln Cys Val Asp Ile Pro Ala Asp
 50 50 55 60
 52 ctg cgg ctg tgc cac aac gtg ggc tac aag aag atg gtg ctg ccc aac 539
 53 Leu Arg Leu Cys His Asn Val Gly Tyr Lys Lys Met Val Leu Pro Asn
 54 65 70 75
 56 ctg ctg gag cac gag acc atg gcg gag gtg aag cag cag gcc agc agc 587
 57 Leu Leu Glu His Glu Thr Met Ala Glu Val Lys Gln Gln Ala Ser Ser
 58 80 85 90 95
 60 tgg gtg ccc ctg ctc aac aag aac tgc cac gcc ggg acc cag gtc ttc 635
 61 Trp Val Pro Leu Leu Asn Lys Asn Cys His Ala Gly Thr Gln Val Phe
 62 100 105 110

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65 Leu Cys Ser Leu Phe Ala Pro Val Cys Leu Asp Arg Pro Ile Tyr Pro
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68 tgt cgc tgg ctc tgc gag gcc gtg cgc gac tcg tgc gag ccg gtc atg      731
69 Cys Arg Trp Leu Cys Glu Ala Val Arg Asp Ser Cys Glu Pro Val Met
70      130      135      140
72 cag ttc ttc ggc ttc tac tgg ccc gag atg ctt aag tgt gac aag ttc      779
73 Gln Phe Phe Gly Phe Tyr Trp Pro Glu Met Leu Lys Cys Asp Lys Phe
74      145      150      155
76 ccg gag ggg gac gtc tgc atc gcc atg acg ccg ccc aat gcc acc gaa      827
77 Pro Glu Gly Asp Val Cys Ile Ala Met Thr Pro Pro Asn Ala Thr Glu
78 160      165      170      175
80 gcc tcc aag ccc caa ggc aca acg gtg tgt cct ccc tgt gac aac gag      875
81 Ala Ser Lys Pro Gln Gly Thr Thr Val Cys Pro Pro Cys Asp Asn Glu
82      180      185      190
84 ttg aaa tct gag gcc atc att gaa cat ctc tgt gcc agc gag ttt gca      923
85 Leu Lys Ser Glu Ala Ile Ile Glu His Leu Cys Ala Ser Glu Phe Ala
86      195      200      205
88 ctg agg atg aaa ata aaa gaa gtg aaa aaa gaa aat ggc gac aag aag      971
89 Leu Arg Met Lys Ile Lys Glu Val Lys Lys Glu Asn Gly Asp Lys Lys
90      210      215      220
92 att gtc ccc aag aag aag aag ccc ctg aag ttg ggg ccc atc aag aag      1019
93 Ile Val Pro Lys Lys Lys Lys Pro Leu Lys Leu Gly Pro Ile Lys Lys
94      225      230      235
96 aag gac ctg aag aag ctt gtg ctg tac ctg aag aat ggg gct gac tgt      1067
97 Lys Asp Leu Lys Lys Leu Val Leu Tyr Leu Lys Asn Gly Ala Asp Cys
98 240      245      250      255
100 ccc tgc cac cag ctg gac aac ctc agc cac cac ttc ctc atc atg ggc      1115
101 Pro Cys His Gln Leu Asp Asn Leu Ser His His Phe Leu Ile Met Gly
102      260      265      270
104 cgc aag gtg aag agc cag tac ttg ctg acg gcc atc cac aag tgg gac      1163
105 Arg Lys Val Lys Ser Gln Tyr Leu Leu Thr Ala Ile His Lys Trp Asp
106      275      280      285
108 aag aaa aac aag gag ttc aaa aac ttc atg aag aaa atg aaa aac cat      1211
109 Lys Lys Asn Lys Glu Phe Lys Asn Phe Met Lys Lys Met Lys Asn His
110      290      295      300
112 gag tgc ccc acc ttt cag tcc gtg ttt aag tga ttctcccggg ggcaggggtg      1264
113 Glu Cys Pro Thr Phe Gln Ser Val Phe Lys
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132 acttacaatt aaggtcaagc ccagaaagtg ataagtgcag ggaggaaaag tgcaagtcca      1804
134 ttatctaata gtgacagcaa agggaccagg ggagaggcat tgctttctct gcccacagtc      1864

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138 tcctatgagc ccggggcatg atctgatccc caagacatgt ggaggggcag cctgtgctg 1984
140 cctttgtgtc agaaaaagga aaccacagtg agcctgagag agacggcgat tttcgggctg 2044
142 agaaggcagt agttttcaaa acacatagtt aaaaaagaaa caaatgaaaa aaatttttaga 2104
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156 gaaacatttc ctttgaactt gattgcctat ggatcaaaga aattcagaac agcctgcctg 2524
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160 ggggtgtccc catccagcga gagagtttca aaagcaaaac atctctgcag tttttcccaa 2644
162 gtaccctgag atacttccca aagcccttat gtttaatcag cgatgtatat aagccagttc 2704
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214 gaaattccc tttcctctag tttcttctct tagtactcct ctttttagatc ctaagtctct 4264
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226 <211> LENGTH: 313
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228 <213> ORGANISM: Homo sapiens
230 <400> SEQUENCE: 2

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232 Met Gly Ile Gly Arg Ser Glu Gly Gly Arg Arg Gly Ala Leu Gly Val
233 1 5 10 15
236 Leu Leu Ala Leu Gly Ala Ala Leu Leu Ala Val Gly Ser Ala Ser Glu
237 20 25 30
240 Tyr Asp Tyr Val Ser Phe Gln Ser Asp Ile Gly Pro Tyr Gln Ser Gly
241 35 40 45
244 Arg Phe Tyr Thr Lys Pro Pro Gln Cys Val Asp Ile Pro Ala Asp Leu
245 50 55 60
248 Arg Leu Cys His Asn Val Gly Tyr Lys Lys Met Val Leu Pro Asn Leu
249 65 70 75 80
252 Leu Glu His Glu Thr Met Ala Glu Val Lys Gln Gln Ala Ser Ser Trp
253 85 90 95
256 Val Pro Leu Leu Asn Lys Asn Cys His Ala Gly Thr Gln Val Phe Leu
257 100 105 110
260 Cys Ser Leu Phe Ala Pro Val Cys Leu Asp Arg Pro Ile Tyr Pro Cys
261 115 120 125
264 Arg Trp Leu Cys Glu Ala Val Arg Asp Ser Cys Glu Pro Val Met Gln
265 130 135 140
268 Phe Phe Gly Phe Tyr Trp Pro Glu Met Leu Lys Cys Asp Lys Phe Pro
269 145 150 155 160
272 Glu Gly Asp Val Cys Ile Ala Met Thr Pro Pro Asn Ala Thr Glu Ala
273 165 170 175
276 Ser Lys Pro Gln Gly Thr Thr Val Cys Pro Pro Cys Asp Asn Glu Leu
277 180 185 190
280 Lys Ser Glu Ala Ile Ile Glu His Leu Cys Ala Ser Glu Phe Ala Leu
281 195 200 205
284 Arg Met Lys Ile Lys Glu Val Lys Lys Glu Asn Gly Asp Lys Lys Ile
285 210 215 220
288 Val Pro Lys Lys Lys Lys Pro Leu Lys Leu Gly Pro Ile Lys Lys Lys
289 225 230 235 240
292 Asp Leu Lys Lys Leu Val Leu Tyr Leu Lys Asn Gly Ala Asp Cys Pro
293 245 250 255
296 Cys His Gln Leu Asp Asn Leu Ser His His Phe Leu Ile Met Gly Arg
297 260 265 270
300 Lys Val Lys Ser Gln Tyr Leu Leu Thr Ala Ile His Lys Trp Asp Lys
301 275 280 285
304 Lys Asn Lys Glu Phe Lys Asn Phe Met Lys Lys Met Lys Asn His Glu
305 290 295 300
308 Cys Pro Thr Phe Gln Ser Val Phe Lys
309 305 310

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VERIFICATION SUMMARY

DATE: 08/26/2004

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11 M:271 C: Current Filing Date differs, Replaced Current Filing Date